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OM nucleic - nucleic search, using sw model

Run on: December 3, 2000, 18:59:08 ; Search time 328.13 Seconds  
(without alignments)  
2843.411 Million cell updates/sec

Title: US-09-227-881-3  
Perfect score: 6169  
Sequence: 1 atctttgttcagtttaccc.....cttgcgcctccatgtcag 6169

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 262060 seqs, 75620727 residues

Total number of hits satisfying chosen parameters: 524120  
Minimum DB seq length: 0  
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%  
Maximum Match 100%

Listing first 45 summaries

Database : Issued\_Patents\_NA:\*

|    |   |
|----|---|
| 1: | /cgn2_6/ptodata/2/1na/5A_COMB.seq:*     |
| 2: | /cgn2_6/ptodata/2/1na/5B_COMB.seq:*     |
| 3: | /cgn2_6/ptodata/2/1na/5C_COMB.seq:*     |
| 4: | /cgn2_6/ptodata/2/1na/5D_COMB.seq:*     |
| 5: | /cgn2_6/ptodata/2/1na/6_COMB.seq:*      |
| 6: | /cgn2_6/ptodata/2/1na/PCITUS_COMB.seq:* |
| 7: | /cgn2_6/ptodata/2/1na/backfiles1.seq:*  |

Pred. No. 19 is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

#### 8

| Result No. | Score | Query | Match | Length | DB                | ID | Description         |
|------------|-------|-------|-------|--------|-------------------|----|---------------------|
| 1          | 640.4 | 10.4  | 1999  | 3      | US-09-645-900A-2  |    | Sequence 2, Appli   |
| 2          | 640.4 | 10.4  | 1999  | 3      | US-09-688-238A-2  |    | Sequence 2, Appli   |
| 3          | 640.4 | 10.4  | 1999  | 3      | US-09-667-790A-2  |    | Sequence 1, Appli   |
| 4          | 638.8 | 10.4  | 1985  | 3      | US-09-791-347-1   |    | Sequence 1, Appli   |
| 5          | 604.4 | 9.8   | 1512  | 3      | US-09-645-900A-3  |    | Sequence 3, Appli   |
| 6          | 604.4 | 9.8   | 1512  | 3      | US-09-682-238A-3  |    | Sequence 3, Appli   |
| 7          | 604.4 | 9.8   | 1512  | 3      | US-09-667-790A-3  |    | Sequence 3, Appli   |
| 8          | 501.8 | 9.8   | 1509  | 3      | US-09-791-347-2   |    | Sequence 2, Appli   |
| 9          | 556.6 | 9.0   | 1968  | 6      | PCT-US95-14024-2  |    | Sequence 2, Appli   |
| 10         | 556.6 | 9.0   | 1969  | 2      | US-09-336-235A-2  |    | Sequence 2, Appli   |
| 11         | 556.6 | 9.0   | 1969  | 2      | US-09-649-432-2   |    | Sequence 2, Appli   |
| 12         | 519.6 | 8.4   | 1491  | 1      | US-09-336-235A-3  |    | Sequence 3, Appli   |
| 13         | 519.6 | 8.4   | 1491  | 2      | US-09-649-432-3   |    | Sequence 3, Appli   |
| 14         | 519.6 | 8.4   | 1491  | 6      | PCT-US95-14024-3  |    | Sequence 3, Appli   |
| 15         | 204   | 3.3   | 643   | 3      | US-09-791-347-5   |    | Sequence 15, Appli  |
| 16         | 174.2 | 2.8   | 282   | 1      | US-09-133-629-8   |    | Sequence 8, Appli   |
| 17         | 173.6 | 2.8   | 17327 | 1      | US-07-906-81-15   |    | Sequence 15, Appli  |
| 18         | 169   | 2.7   | 5543  | 4      | US-09-688-080-01  |    | Sequence 101, Appli |
| 19         | 165   | 2.7   | 2115  | 2      | US-09-395-800A-7  |    | Sequence 7, Appli   |
| 20         | 164.6 | 2.7   | 6769  | 2      | US-09-480-784-20  |    | Sequence 20, Appli  |
| 21         | 164.6 | 2.7   | 6769  | 2      | US-09-483-553-20  |    | Sequence 20, Appli  |
| 22         | 164.6 | 2.7   | 6769  | 2      | US-09-487-002-20  |    | Sequence 20, Appli  |
| 23         | 164.6 | 2.7   | 6769  | 2      | US-09-483-554B-20 |    | Sequence 20, Appli  |
| 24         | 164.6 | 2.7   | 6769  | 2      | US-09-488-011B-20 |    | Sequence 20, Appli  |
| 25         | 164.6 | 2.7   | 6769  | 6      | PCT-US95-10202-20 |    | Sequence 20, Appli  |
| 26         | 164.6 | 2.7   | 6769  | 6      | PCT-US95-10203-20 |    | Sequence 20, Appli  |

| ALIGNMENTS                                       |       |
|--|-------|
| RESULT   | 1     |
| US-08-645-900A-2                                 |       |
| Sequence 2, Application US/08645900A             |       |
| PATENT NO. 5849879                               |       |
| GENERAL INFORMATION:                             |       |
| APPLICANT: NGUYEN, THAI D.                       |       |
| APPLICANT: POLANSKY, JON R.                      |       |
| APPLICANT: HUANG, WIDONG                         |       |
| TITLE OF INVENTION: METHODS FOR THE DIAGNOSIS OF |       |
| TITLE OF INVENTION: GLAUCOMA                     |       |
| NUMBER OF SEQUENCES: 3                           |       |
| CORRESPONDENCE ADDRESS:                          |       |
| ADDRESSEE: HOWREY & SIMON                        |       |
| STREET: 129 PENNSYLVANIA AVE., N.W.              |       |
| CITY: WASHINGTON                                 |       |
| STATE: D.C.                                      |       |
| COUNTRY: US                                      |       |
| ZIP: 20004                                       |       |
| COMPUTER READABLE FORM:                          |       |
| MEDIUM TYPE: FLOPPY DISK                         |       |
| COMPUTER: IBM PC COMPATIBLE                      |       |
| OPERATING SYSTEM: PC-DOS/MS-DOS                  |       |
| SOFTWARE: Patent In Release #1.0, Version #1.25  |       |
| CURRENT APPLICATION DATA:                        |       |
| APPLICATION NUMBER: US/08/645, 900A              |       |
| FLING DATE: 14 MAY 1996                          |       |
| CLASSIFICATION: 530                              |       |
| ATTORNEY/AGENT INFORMATION:                      |       |
| NAME: MARSH, DAVID                               |       |
| REGISTRATION NUMBER: 41,408                      |       |
| TELECOMMUNICATION INFORMATION:                   |       |
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| TELEFAX: (202) 383-6610                          |       |
| INFORMATION FOR SEQ ID NO: 2:                    |       |
| SEQUENCE CHARACTERISTICS:                        |       |
| LENGTH: 1999 base pairs                          |       |
| TYPE: nucleic acid                               |       |
| STRANDEDNESS: single                             |       |
| TOPOLOGY: linear                                 |       |
| MOLECULE TYPE: cDNA                              |       |
| US-08-645-900A-2                                 |       |
| Query Match                                      | 10.4% |
| Best Local Similarity                            | 99.8% |
| Matches  | 641   |
| Matches 641; Conservative                        | 0     |
| Matches 641;保守                                   | 0     |
| Matches  | 0     |
| Indels   | 0     |
| Gaps   | 0     |
| Gaps 0   | 0     |
| Sequence 5, Appli                                |       |
| Sequence 20, Appli                               |       |
| Sequence 5, Appli                                |       |
| Sequence 28, Appli                               |       |
| Sequence 5, Appli                                |       |
| Sequence 5, Appli                                |       |
| Sequence 5, Appli                                |       |
| Sequence 2, Appli                                |       |
| Sequence 3, Appli                                |       |
| Sequence 1, Appli                                |       |
| Sequence 1, Appli                                |       |
| Sequence 7, Appli                                |       |
| Sequence 7, Appli                                |       |
| Sequence 7, Appli                                |       |
| Sequence 20, Appli                               |       |
| Sequence 20, Appli                               |       |
| Sequence 22, Appli                               |       |
| Sequence 1, Appli                                |       |
| Sequence 17, Appli                               |       |















OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/649,432  
FLILING DATE: 17 MAY 1996  
CLASSIFICATION: 435  
PRIORITY APPLICATION DATA:  
APPLICATION NUMBER: 08/336,235  
FLILING DATE:  
ATTORNEY/AGENT INFORMATION:  
NAME: AUERBACH, JEFFREY I  
REGISTRATION NUMBER: 32,780  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (202) 383-7451  
TELEFAX: (202) 383-6010  
SEQUENCE INFORMATION FOR SEQ ID NO: 3:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1491 base pairs  
TYPE: nucleic acid  
STRANDEDNESS: double  
TOPOLOGY: linear  
MOLECULE TYPE: cDNA  
HYPOTHETICAL: NO  
ANTI-SENSE: NO  
IMMEDIATE SOURCE:  
CLONE: TIGR coding sequence



|    |      |  |       |
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| Db | 1    |  | 60    |
| Oy | 6001 | ctacaggcgctccaaagcctc---cctgcctttctctcgagactgcagactagcaca      | 6056C |
| Db | 61   |  | 120   |
| Oy | 6057 | agacatgtgatattaaaggaagcaca-cgcatacacttcataattttacttagaaatggatc | 6115  |
| Db | 121  |  | 180   |
| Oy | 6116 | AGACAGATGAAATTAGGAAGACACAGCGATCACCTCAAGTATTACTGAAATTAGCTC      | 6169  |
| Db | 181  |  | 234   |
|    |      | ctgagggctatgtgatagtggttcagttctgtgcggccatcag                    |       |
|    |      | CTGAGAGCTCATTTAGATGATGTGTCAGGTCTGTGTCGCCCTCCATGTCAG            |       |

Search completed: December 3, 2000, 19:00:01  
Job time: 41727 sec

